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J Schultz, F Milpetz, P Bork, CP ... - Proceedings of the ..., 1998 - National Acad Sciences  
... Recent reports (43, 44) **identify** the rostral cerebellar malformation gene product (rcm) and similar homologues ... As the DEATH domain of UNC-5 is not **annotated** in databases, this is ... in cell signaling (7). In accordance with this, the SMART alignment **database** contains several ...

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SG Megason, SE Fraser - Mechanisms of development, 2003 - Elsevier  
... image sets and manage the large amount of data through the use of a **database**. ... all these dimensions with sufficient spatial and temporal resolution for tracking **cells** is a ... Principle among these is achieving sufficient spatial and temporal resolution for **identifying** and tracking **cells** ...

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L Malmstrom, J Malmstrom, G Marko-Varga, G ... - Journal of Proteome ..., 2002 - pubs.acs.org  
... Hence, **identifying** and **filtering** interesting data is necessary because of the large amount ... Our research team uses the **database** to investigate new hypothesis within human pulmonary ... Today, the 2DDB retrieves information about **annotated** proteins from Internet databases such ...

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X Zhou, MCJ Kao, WH Wong - Proceedings of the National ..., 2002 - National Acad Sciences  
... We check the GO process classification of the transitive genes on the SP to **identify** the L0 and L1 matches, as discussed in ... The 51 novel genes predicted with the conservative rule for which no biological process **annotation** is available in either **database** are listed in ...

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[MIPS Arabidopsis thaliana Database \(MatDB\): an integrated biological knowledge ...](#) - [nih.gov](#)

H Schoof, P Zaccaria, H Gundlach, K Lemcke, S ... - Nucleic acids ..., 2002 - Oxford Univ Press  
... Thus, manual **annotation** by experts for certain protein families should be available in a whole-genome **database**. ... structure of the phenotype catalog was designed to aid researchers observing a phenotype to query the **database** and **identify** characterized mutants with a ...

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V Brusic, J Zelezniukow, N Petrovsky - Journal of immunological methods, 2000 - Elsevier  
... the blind panel study were subjected to statistical analysis to **identify** possible antibody ... This is an example of a specialised **database** providing information related to a ... Users can extract fully **annotated** sequence information, pairwise comparisons, multiple sequence alignments ...

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[Database tools for integrating and searching membrane property data correlated ...](#)

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... to put only positive data in the **database** (data which affirms the presence of a property in a cell

compartment). ... Thus, gatekeeping is kept to a minimum; the functions of the **database** managers are to ensure that the data are formatted properly and **annotated** appropriately. ...

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[PDF] ► [Helping data suppliers and consumers negotiate the details: a \*\*database\*\* view ...](#)

A Rosenthal, E Scors - submitted to Conference on Cooperative Information ..., 1999 - Citeseer

... We now try to capture the requirements implied by the above scenarios, and to **identify** the major functions of a tool that might meet them. ... We assume there is an **annotation** facility for attaching information to various granules of a **database** (eg, a table, column, **cell**, or ...

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[Models and tools for collaborative \*\*annotation\*\*](#) - ► [arxiv.org](#) [PDF]

X Ma, H Lee, S Bird, K Masda - Arxiv preprint cs/0204004, 2002 - arxiv.org

... and **identify** particular vocalizations of interest, noting their start and end times and classifying the call types. ... The following table shows statistics for  $K \times$  for the test **database**. ... Consider a boolean  $n \times n$  matrix, where  $n$  is the number of anchors in an **annotation** graph. ...

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[PDBsum: a Web-based \*\*database\*\* of summaries and analyses of all PDB structures](#)

RA Laskowski, EG Hutchinson, AD Michie, AC ... - Trends in Biochemical ..., 1997 - Elsevier

... our own CATH classification of protein structural families and the EC, PDB Enzyme Structures

**Database**. ... by the SITE records in the PDB file, while the small red dots **identify** the residues ... 3).

Further details on all the **annotations** are provided by the links (not shown here) that ...

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